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Result
No.
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N_Geneseq_1101:*
1: /SIDS2/gcgdata
2: /SIDS2/gcgdata
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Match Length
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Rev4 coding sequen PCRIL DNA coding sequen PCRIL DNA coding sequent pro-magainin codin Human breast and o Arabidopsis thalia Arabidopsis thalia
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ALIGNMENTS

AAA28510 RESULT

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AAA28510 standard; DNA; 39 BP.

AAA28510;

29-AUG-2000

(first entry)

Rev4 coding sequence

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Polypeptides for reducing proteolytic degradation of proteins administered to, or produced by a plant comprise indolicin or its
                             WPI; 2000-365597/31.
P-PSDB; AAY92796.
                                                                                                          30-OCT-1998;
02-NOV-1998;
                                                         Everett NP, Li Q,
                                                                            (INTE-) INTERLINK BIOTECHNOLOGIES LLC. (KENT ) UNIV KENTUCKY RES FOUND.
                                                                                                                                                         11-MAY-2000.
                                                                                                                                                                             WO200026344-A1.
                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                Magainin; antimicrobial; transgenic plant; protease degradation; Rev4; indolicidin; protein production; reverse peptide; ss.
                                                                                                                                      29-OCT-1999;
                                                                                                          98US-0106373.
98US-0106537.
                                                                                                                                      99WO-US25561
                                                          Lawrence C,
                                                           Davies
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Best Local S
Matches 39
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             Indolicidin is a potent antimicrobial tridecapeptide, originally purified from cytoplasmic granules of bovine neutrophils. Reverse peptide, Rev4 of indolicidin (see AAY92794) was found to have increased stability against plant protease degradation. Expression of antimicrobial stability against plant protease degradation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCRIL DNA coding sequence
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                                                                                                                               Polypeptides for reducing proteolytic degradation of proteins administered to, or produced by a plant comprise indolicin or
                                                                                                                                                                                                                                                                                           30-OCT-1998;
02-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                      Example 17; Page 35; 50pp; English.
                                                                                                                                                                                                                      Everett NP,
                                                                                                                                                                                                                                                                                                                                       29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Magainin; antimicrobial; transgenic plant; protease degradation; Rev4;
                                                                                                                 functional equivalents
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39; Conservative
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transgenic plants suffers a major limitation
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98US-0106537.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      production; reverse peptide; ss.
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Pred.
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3.8e-06;
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               antimicrobial
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RESULT 3
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AAA28512-13 are primers used to introduce a secretion signal peptide
from tobacco PR-1b to RIL (encoding Rev4) to facilitate secretion of Rev4
cinto the extracellular space of plant tissues where invading pathogens
might be first encountered. Indolicidin is a potent antimicrobial
tridecapeptide, originally purified from cytoplasmic granules of bowine
neutrophils. Reverse peptide, Rev4 of indolicidin (see AAY92794) was
found to have increased stability against plant protease degradation.
Expression of antimicrobial peptides in transgenic plants suffers a
major limitation in that the foreign peptides are susceptible to rapid
degradation by proteases. The invention concerns reducing the extent of
protease degradation of a protein applied to, or produced by a plant by
administering indolicidin, Rev4 or a functional equivalent to the plant.
Transgenic plants expressing indolicidin and Rev4 are useful for
production of the antimicrobial peptides. Compositions containing
indolicidin and Rev4 are also useful for production of agronomically
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Best Local
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                                                                                                                                                                                                                                                                                                                                        Polypeptides for reducing administered to, or produc
                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-365597/31.
                                                                                                                                                                                                                                                                                                                                                                                                                               Everett NP, Li Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-OCT-1998;
02-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200026344-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nicotiana tabacum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer RIL 3' SacI for tobacco PR-lb signal peptide coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-2000 (first entry)
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                                                                                                                                                                                                                                                                                      Example 15; Page 35; 50pp; English.
                                                                                                                                                                                                                                                                                                                     functional equivalents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Magainin; antimicrobial; transgenic plant; protease degradation; Rev4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INTE-) INTERLINK BIOTECHNOLOGIES LLC. (KENT ) UNIV KENTUCKY RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               indolicidin; protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 49 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0106373.
98US-0106537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US25561.
                                                                                                                                                                                                                                                                                                                                         ducing proteolytic de produced by a plant
                                                                                                                                                                                                                                                                                                                                                                                                                                 Lawrence C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50 C; 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 Davies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 72 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reverse peptide;
                                                                                                                                                                                                                                                                                                                                           degradation of proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                 ΜĦ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.4e-06;
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important proteins in plants

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RESULT 4
AAA28514/c
ID AAA285
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   Query Match
Best Local Similarity
Matches 37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 52 BP; 15 A; 16 C; 11 G; 10 T; 0 other;
                                                                                                                     increased stability against plant protease degradation. Expression of antimicrobial peptides in transgenic plants suffers a major limitation in that the foreign peptides are susceptible to rapid degradation by proteases. The invention concerns reducing the extent of protease degradation of a protein applied to, or produced by a plant by administering indolicidin, Rev4 or a functional equivalent to the plant. Transgenic plants expressing indolicidin and Rev4 are useful for production of the antimicrobial peptides. Compositions containing indolicidin and Rev4 are also useful for production of agronomically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA28514;
                                                                                                                                                                                                                                                          A gene (Pre-Pro-RIL) encoding a reverse indolicidin peptide, Rev4, for secretion in plants was constructed using a pro-sequence based on that found in natural magainin genes. The pro-sequence was linked to Rev4 through a His-Ser motif that corresponds to the site at which plant extracellular proteases cleave natural magainins. Rev4 was found to have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200026344-A1
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                                                                            Sequence 52
                                                                                                                                                                                                                                                                                                                                                     Example 16; Page 35; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                            Polypeptides for reducing proteolytic degradation of proteins administered to, or produced by a plant comprise indollcin or
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-365597/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KENT ) UNIV KENTUCKY RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INTE-) INTERLINK BIOTECHNOLOGIES LLC.
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02-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rana pipiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              indolicidin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Magainin; antimicrobial; transgenic plant; protease degradation; Rev4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pro-magainin coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                 functional equivalents
                                                                                                          important proteins in plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 ggagatggccttggtggccttggaaatggcctcttatt 39
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Conservative 0
                                                                              BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li Q,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein production; reverse peptide; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0106373.
98US-0106537.
                                                                               A; 14 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lawrence C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97.4%;
                                94.98;
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Pred. No.
       0;
                  Score 37; ; Pred. No.
                                                                              10 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Davies MH
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9.8e-06;
                    DB 21; 1
2.4e-05;
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                                  Length 52;
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         Indels
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         0;
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         Gaps
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RESULT
AAF21668/
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B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Addison's disease; allergy; autoimmune haemolytic anaemia; autoimmune thyroiditis; diabetes mellitus; Crohn's disease; multiple sclerosis; rheumatoid arthritis; ulcerative coliti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nootropic; neurpprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human breast and ovarian cancer associated antigen gene SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; breast cancer; ovarian cancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-2001
                                                                                                                                                                                            nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic and cardiant activity. The polynucleotide and protein sequences are used in the diagnosis of cancer,
                                                                                                                                                                                                                                                                                                                                    Sequences AAF21614 - AAF22031 represent DNA sequences encoding human proteins AAB58711 - AAB59128. The DNA and protein sequences are associated with breast and ovarian cancer. Included in the invention are sequences AAF22032 - AAF22040 and AAB59129 which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human breast and ovarian cancer associated gene sequences and the polypeptides encoded by these genes, useful in the prevention, treatment and diagnosis of cancer, immune disorders, cardiovascular disorders and neurological diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cardiovascular disorder; wound healing; neurological disease;
                                                       diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; cardiovascular disorders such as myocardial ischaemias; wound healing; neurological diseases such as
                                                                                                                 particularly breast and ovarian cancer. The nucleic acid sequences, proteins, agonists and agonists may also be used in the diagnosis, prevention and treatment of immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis,
                                                                                                                                                                                                                                                                               invention. The breast and ovarian cancer associated DNA, protein, agonist or antagonist sequences exhibit cytostatic; immunosuppressive;
                                                                                                                                                                                                                                                                                                              isolation and characterisation of the DNA and protein sequences of the
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 520; 1299pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P-PSDB; AAB58765.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200055173-A1
                                         cerebral anoxia and epilepsy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  52 GAGATGGCCTTGGTGGCCTTGGAAATGGCCTCTTATT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0124270
                                           and infectious diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          arthritis; ulcerative colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunosuppressive;
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994 BP;

217 A;

298 C;

290 G; 184 T; 5 other;

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	16-APR-1999; 19-APR-1999; 21-APR-1999; 23-APR-1999; 23-APR-1999; 23-APR-1999; 30-APR-1999; 30-APR-1999; 30-APR-1999; 30-APR-1999; 10-MAY-1999; 11-MAY-1999;	• ਜ	RESULT 6 AAC44240 stan AC AAC44240 stan XX AAC44240; XX DT 18-OCT-2000 XX XX Arabidopsis t XX W Hybridisation KW Protein ident KW protein ident KW protein ident KW protein ident	Best Local Simil Matches 26; C 5 gatggcct 432 GATGGCCA
99US-0135299. 99US-0136021. 99US-0136021. 99US-013692. 99US-0137222. 99US-0137722. 99US-0137724. 99US-013762. 99US-0138094. 99US-0138094. 99US-0138094. 99US-0138094. 99US-0138094. 99US-0138094.	99US-012845. 99US-0130077. 99US-0130449. 99US-0130449. 99US-0130891. 99US-0131449. 99US-0132407. 99US-0132486. 99US-0132486. 99US-0132486. 99US-0132486. 99US-0132486. 99US-0132486. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134286. 99US-0134218. 99US-0134218. 99US-0134218. 99US-0134370. 99US-0134370.	P-0301439. P-0301439. S-0121825. S-0123548. S-0125788. S-0126785. S-0126785. S-0127462. S-0127462.	AAC44240 standard; DNA; 315 BP. AAC44240; 18-OCT-2000 (first entry) Arabidopsis thaliana DNA fragment SEQ ID NO: 42138. Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.	pal Similarity 81.2%; Pred. No. 17; 26; Conservative 0; Mismatches 6; Indels 0; Gaps gatggccttggtggccttggaaatggcctctt 36
				0;
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990S-01463 990S-01463 990S-01472 990S-01472 990S-01473 990S-01473 990S-01473 990S-01474 990S-01474 990S-01474 990S-01474 990S-01474 990S-01474 990S-01474	99US-01443334 99US-01443354 99US-01444355 99US-0144884 99US-0145086 99US-0145086 99US-0145087 99US-0145087 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145 99US-0145145	990S-0141847 990S-0141847 990S-0142154 990S-0142154 990S-0142803 990S-0142803 990S-0142803 990S-0142803 990S-014297 990S-014297 990S-0144085 990S-0144085 990S-0144086 990S-0144325 990S-0144333	99US-0139 99US-0139 99US-0139 99US-0139 99US-0139 99US-0139 99US-0139 99US-0139 99US-0139 99US-0140 99US-0140	99US-0139452 99US-0139453 99US-0139453 99US-0139454 99US-0139455 99US-0139455

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אם קרים את המוצא המוצא
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                                                                                                                      Query Match
Best Local S
Matches 25
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13-AUG-1999;
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17-AUG-1999;
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15-SEP-1999;
16-SEP-1999;
20-SEP-1999;
22-SEP-1999;
23-SEP-1999;
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24-SEP-1999;
26-OCT-1999;
26-OCT-1999;
27-OCT-1999;
28-OCT-1999;
      203 aggagctggtcttggtgggcttgggattgg 232
                                    1 aggagatggccttggtggccttggaaatgg 30
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25; Conservative
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990S-0148368.
990S-0149368.
990S-0149723.
990S-0149929.
990S-0149929.
990S-0150884.
990S-0151065.
990S-0151065.
990S-0151080.
990S-015928.
990S-015928.
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990S-016076.
990S-0161406.
990S-0161406.
990S-016136.
990S-016136.
990S-016136.
990S-016136.
990S-016136.
990S-016136.
                                                                                                                                                       56.4%;
                                                                                                                            0;
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Pred. No. 19;
0; Mismatches
                                                                                                                                                                                           21;
                                                                                                                                  5
                                                                                                                                                                                        Length 315;
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                                                                                                                                     0;
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   RESULT
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16-JUN-1999;
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   990S-0121825.
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990S-014254
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990S-0145218
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990S-0145228
RESULT
AAC44352
ID AAC4
XX
AC AAC4
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Best Local s
Matches 25
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23-AUG-1999
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26-AUG-1999
27-AUG-1999
28-EP-1999
28-EP-1999
29-SEP-1999
21-CUT-1999
22-CUT-1999
23-CUT-1999
                                  AAC44352 standard; DNA; 506
           AAC44352;
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                                                                                                                                               Similarity 83.3
25; Conservative
                                                                                                                                                                                              9908-0149723

9908-0149902

9908-0149902

9908-0150884

9908-0151086

9908-0151086

9908-0151080

9908-01511303

9908-0151303

9908-0153758

9908-0153758

9908-0153758

9908-0155479

9908-0155486

9908-0155486

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9908-0161361
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                                ВP
                                                                                                                                               0;
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Pred. No. 19;
0; Mismatches
                                                                                              230
                                                                                                                      30
                                                                                                                                                                      21;
                                                                                                                                             <u>ა</u>
                                                                                                                                                                    Length 322;
                                                                                                                                             Indels
                                                                                                                                            0;
                                                                                                                                           Gaps
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18-JUN-1999
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18-JUN-1999 22-JUN-1999 23-JUN-1999 24-JUN-1999 24-JUN-1999 25-JUN-1999 26-JUN-1999 27-JUN-1999 28-JUN-1999 29-JUN-1999 20-JUL-1999 20-JUL-1999 21-JUL-1999 21-JUL
99US-0139763 99US-0139817 99US-0140354 99US-0140823 99US-0141287 99US-0142055 99US-0142205 99US-0142320 99US-0143542 99US-0143542 99US-0144085 99US-0144085 99US-0144333 99US-0144334 99US-0144334 99US-0145085 99US-0145218 99US-0145218 99US-0145218 99US-0145218 99US-0145218 99US-0145218 99US-0145386 99US-0147303 99US-0149356 99US-0149368 99US-0149329 99US-0149329 99US-0149368 99US-0149329 99US-0149330 99US-0149330 99US-0149330 99US-0149336

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RESULT
AAC40689
ID AAC4
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XC AAC4
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Best Local Similarity
Matches 25; Conserv
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31-AUG-1999
01-SEP-1999
07-SEP-1999
15-SEP-1999
15-SEP-1999
22-SEP-1999
22-SEP-1999
23-SEP-1999
24-SEP-1999
25-OCT-1999
06-OCT-1999
07-OCT-1999
07-OCT-1999
13-OCT-1999
 Hybridisation assay; genetic mapping; gene expression protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.
                                       Arabidopsis
                                                          17-OCT-2000
                                                                             AAC40689;
                                                                                              AAC40689 standard;
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                                                                                                                                          l aggagatggccttggtggccttggaaatgg
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20 aggagctggtcttggtgggcttgggattgg
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                                      thaliana DNA fragment SEQ ID NO: 29206.
                                                                                                                                                                                  Conservative
                                                         (first
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9908-0151080
9908-0151303
9908-0151393
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9908-0155486
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                                                                                              DNA;
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                                                                                                                                                                                Score 22; DB Pred. No. 21; 0; Mismatches
                                                                                              ВP
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                    control;
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 25-MEB-1999
05-MAR-1999
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29-MAR-1999
29-MAR-1999
16-APR-1999
06-APR-1999
08-APR-1999
11-APR-1999
23-APR-1999
23-APR-1999
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28-APR-1999
29-MAY-1999
21-MAY-1999
21-JUN-1999
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RESULT 1
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Best Local S
Matches 25
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25-SEP-1999
26-OCT-1999
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21-OCT-1999
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                                                               Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.
        06-SEP-2000
                                                                                                                                18-OCT-2000 (first entry)
                            EP1033405-A2
                                               Arabidopsis thaliana.
                                                                                                          Arabidopsis thaliana DNA fragment SEQ ID NO: 70502.
                                                                                                                                                        AAC52162;
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00 aggagctggtcttggtgggcttgggattgg 22:
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25; Conservative
                                                                                                                                                                           standard; DNA; 556 BP.
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99US-0154039

99US-0155486

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99US-0156596

99US-0156596

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                                                                                                                                                                                                                                                                                          21;
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                                                                                                                                                                                                                                                                     Gaps
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29-JUN-1999 30-JUL-1999 01-JUL-1999 06-JUL-1999 08-JUL-1999 11-JUL-1999 12-JUL-1999 13-JUL-1999 13-JU

99US-0140991 99US-0141287 99US-014287 99US-0142390 99US-0142390 99US-0142390 99US-0142970 99US-0144005 99US-0144005 99US-0144005 99US-0144005 99US-0144005 99US-0144086 99US-014433 99US-014433 99US-014433 99US-014433 99US-014433 99US-014433 99US-014433 99US-014433 99US-014433 99US-014591 99US-014591 99US-014591 99US-014591 99US-014703 99US-014703 99US-014703 99US-014703 99US-014703 99US-014703 99US-014972 99US-014930 99US-014930 99US-014930 99US-014930 99US-014930 99US-0151066 99US-0151066 99US-0151066 99US-0151066 99US-0151033 99US-0151333 99US-0151333 99US-0151333 99US-0151333

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99US-0139458 99US-0139459 99US-0139461 99US-0139463 99US-0139750 99US-0139763 99US-0139763 99US-0139899 99US-0140353 99US-0140823 99US-0140823 99US-0141287 99US-0141287 99US-0141287 99US-0141287 99US-0141287 99US-0141287 99US-0142390 99US-0142390	99US-0135629. 99US-0136392. 99US-0136782. 99US-0137528. 99US-0137502. 99US-0137502. 99US-0138094. 99US-0138540. 99US-0138847. 99US-0138847. 99US-0139452. 99US-0139453. 99US-0139454. 99US-0139454. 99US-0139456.	990S-0131449. 990S-0131449. 990S-0132407. 990S-0132485. 990S-0132486. 990S-0132486. 990S-0132487. 990S-0134218. 990S-0134218. 990S-013421. 990S-013470. 990S-0134370. 990S-0134370. 990S-0134370. 990S-0134370. 990S-0134370.	2000EP-0301439. 99US-0121825. 99US-0123180. 99US-0125788. 99US-0126764. 99US-0126785. 99US-0126785. 99US-0126785. 99US-012834. 99US-0130670. 99US-0130610.
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RESULT 1
AAC40427
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Matches 25
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29-SEP-1999
04-CCT-1999
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17-CCT-1999
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29-CCT-1999
25-FEB-1999;
05-MAR-1999;
09-MAR-1999;
23-MAR-1999;
25-MAR-1999;
                                                                                                                                    Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.
                                                                                                                  Arabidopsis thaliana.
                                                                                                                                                                           Arabidopsis thaliana DNA fragment SEQ ID
                                                                                                                                                                                                                                    AAC40427 standard; DNA; 585
                                                         25-FEB-2000; 2000EP-0301439
                                                                            06-SEP-2000.
                                                                                               EP1033405-A2
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                                                                                                                                                                                                                  AAC40427;
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25; Conservative
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990S-0159331

990S-0159637

990S-015968

990S-015968

990S-0160761

990S-0160770

990S-0160770

990S-0160815

990S-0160815

990S-0160816

990S-0160981

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990S-0161406

990S-0161361

990S-0161361

990S-0161361

990S-0161932

990S-0161932

990S-0161932

990S-0161932
99US-0121825.
99US-0123180.
99US-0123548.
99US-0125788.
99US-0126264.
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99US-0156596.
99US-0157753.
99US-0157753.
99US-0158029.
99US-0158232.
99US-0158369.
99US-0158369.
99US-0159293.
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                                                                                                                                                                                                                                                                                                                         ; Score 22; DB;
; Pred. No. 21;
0; Mismatches
                                                                                                                                                                                                                                    BP.
                                                                                                                                                                            NO: 28267.
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l Similarity 83.3%;
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Similarity 83.3%;
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RESULT :
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Best Local 9
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05-MAR-1999

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29-OCT-1999;
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Best Local Similarity 83.3
Matches 25; Conservative
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||||| ||| ||| |||||| |||| ||| |||
203 aggagctggtcttggtgggcttgggattgg 232
                                                                                                                                                                            56.4%; Score 22; DB 21; Length 775; B3.3%; Pred. No. 23; tive 0; Mismatches 5; Indels
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